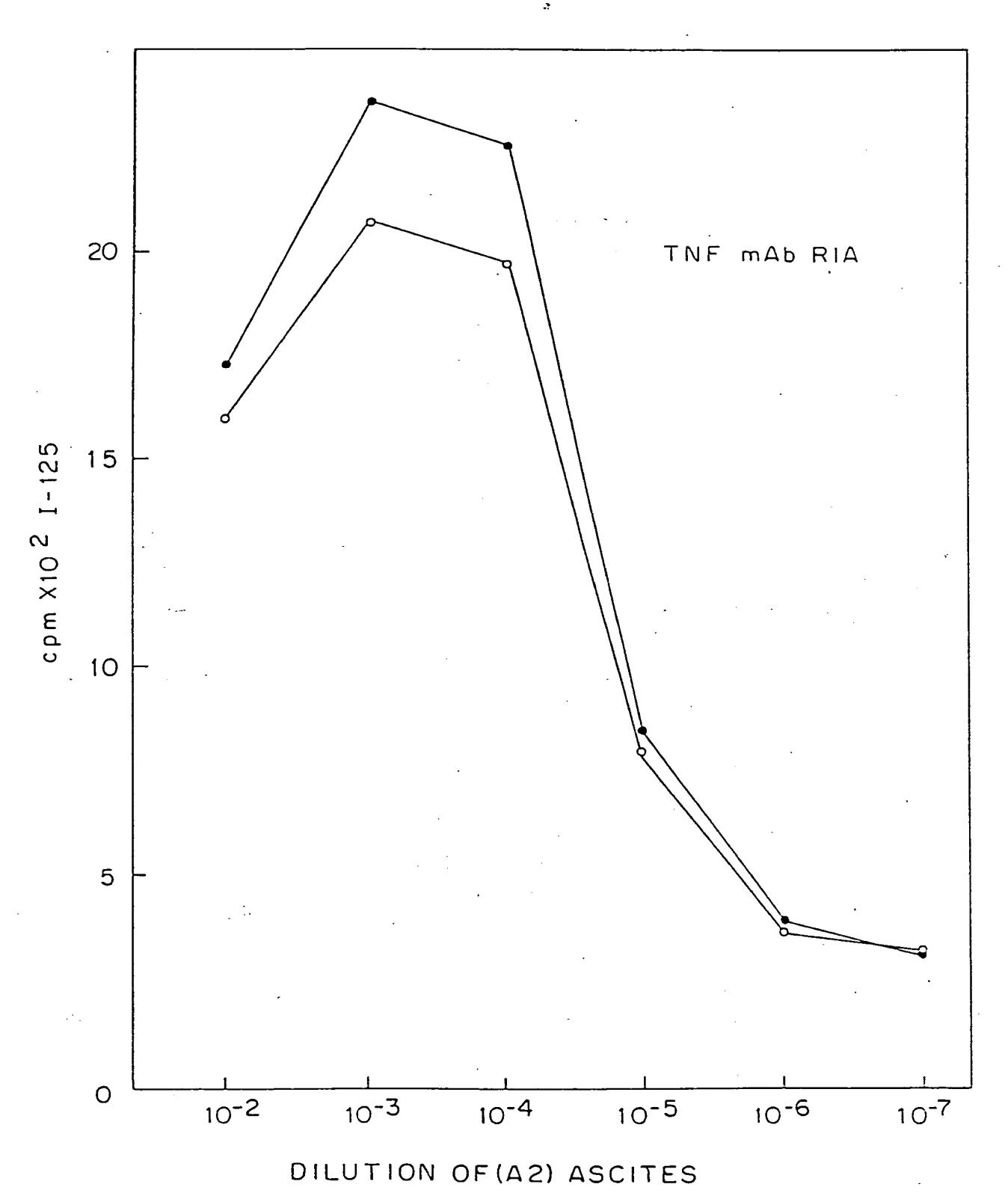
FIG. 1



F I G₂. 2

27:5

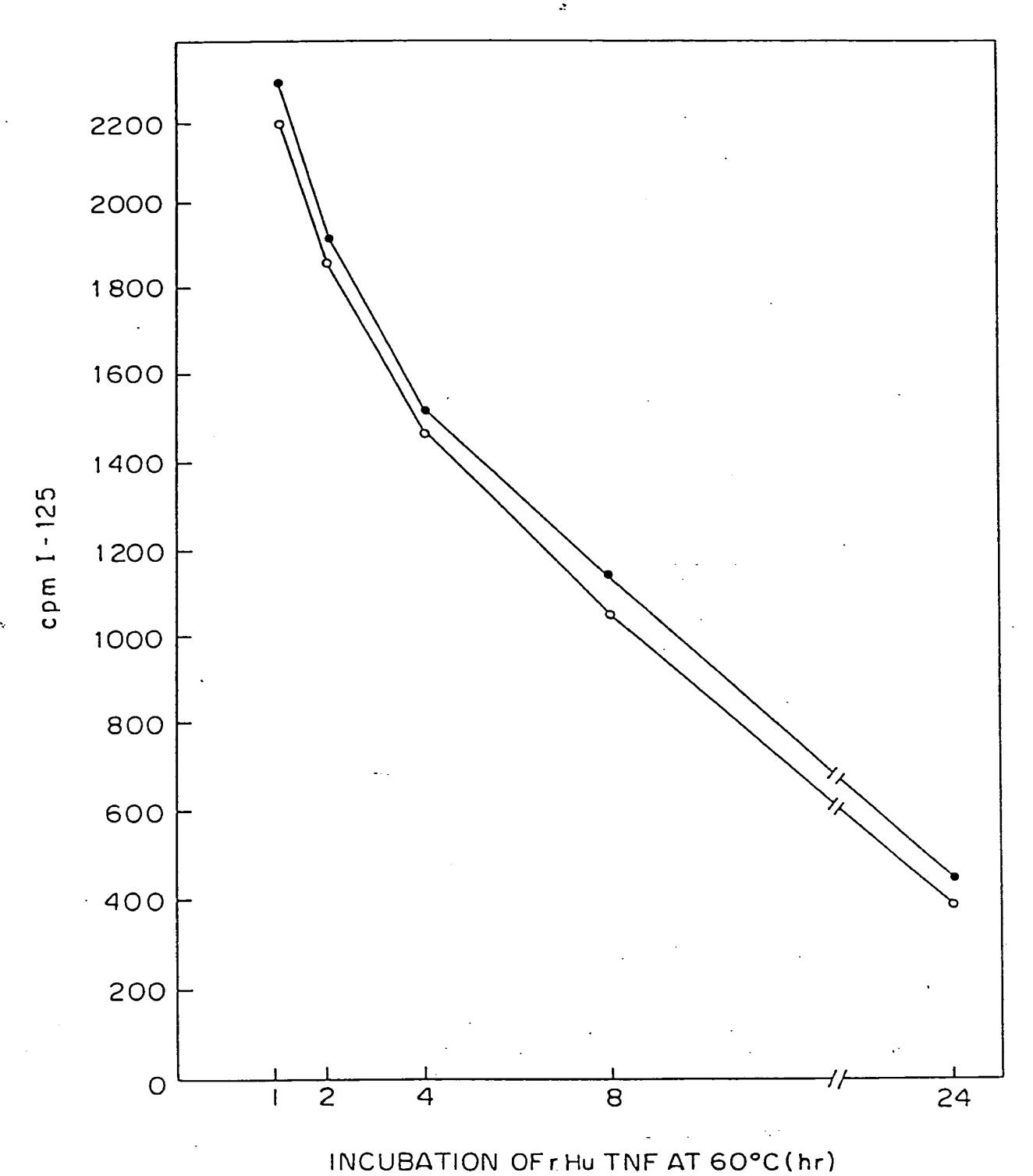
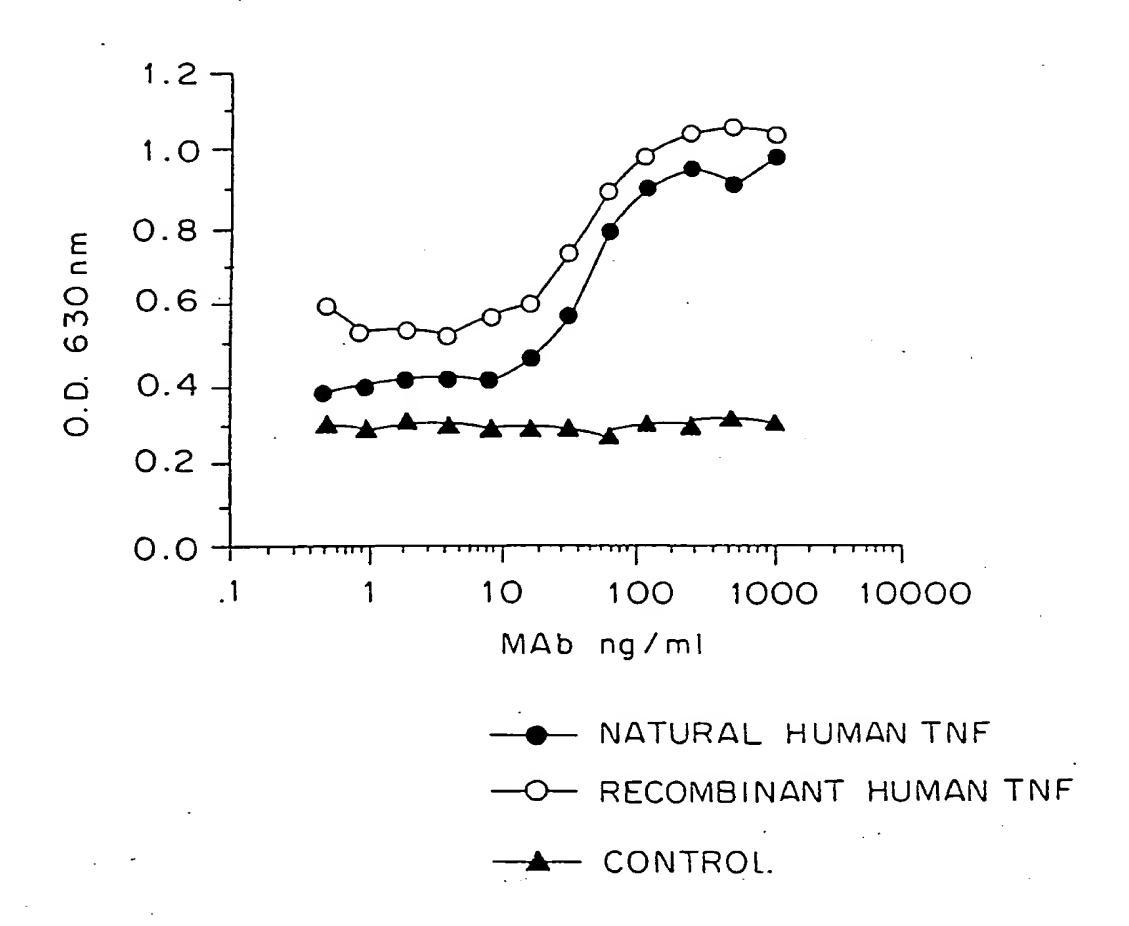
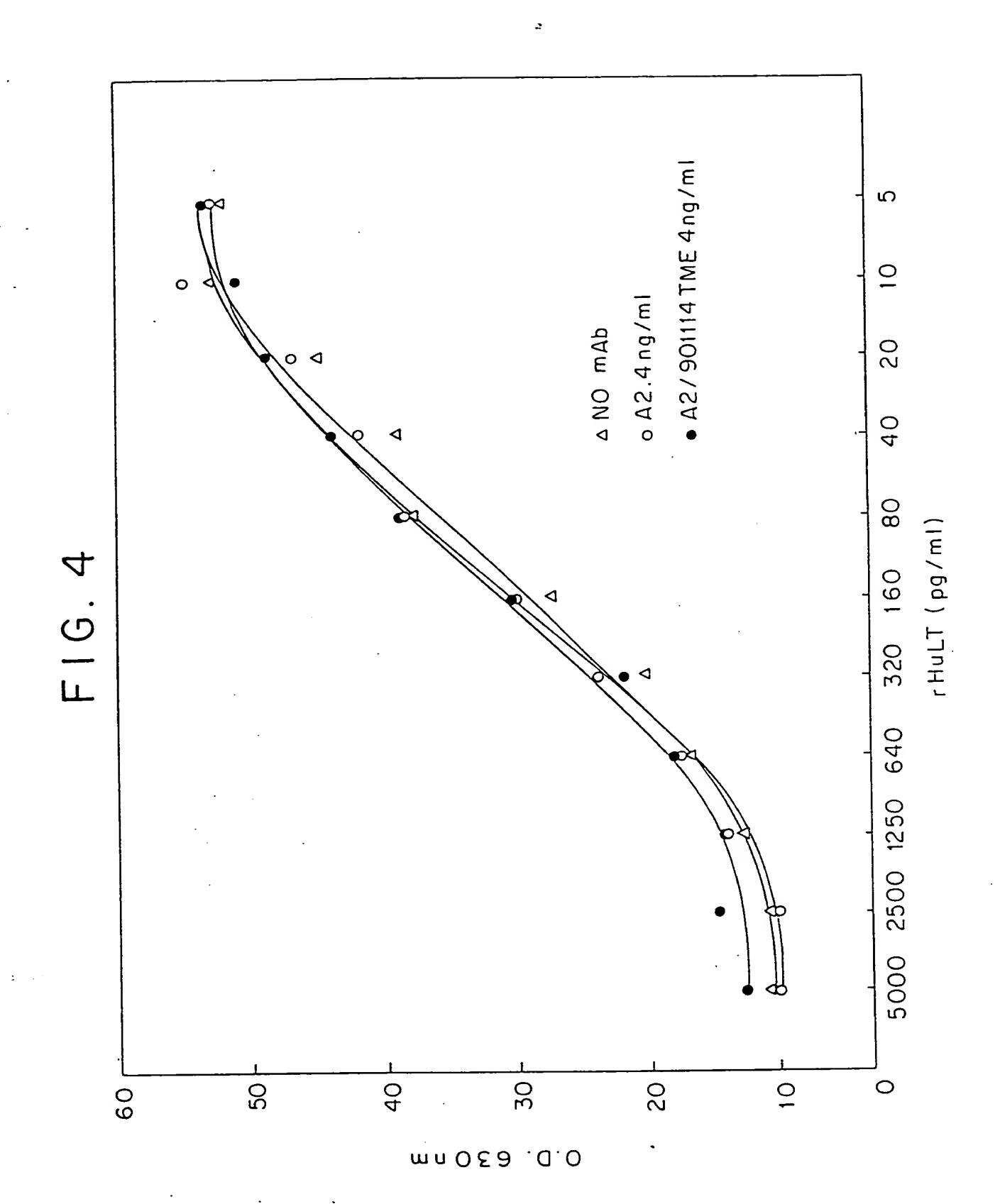
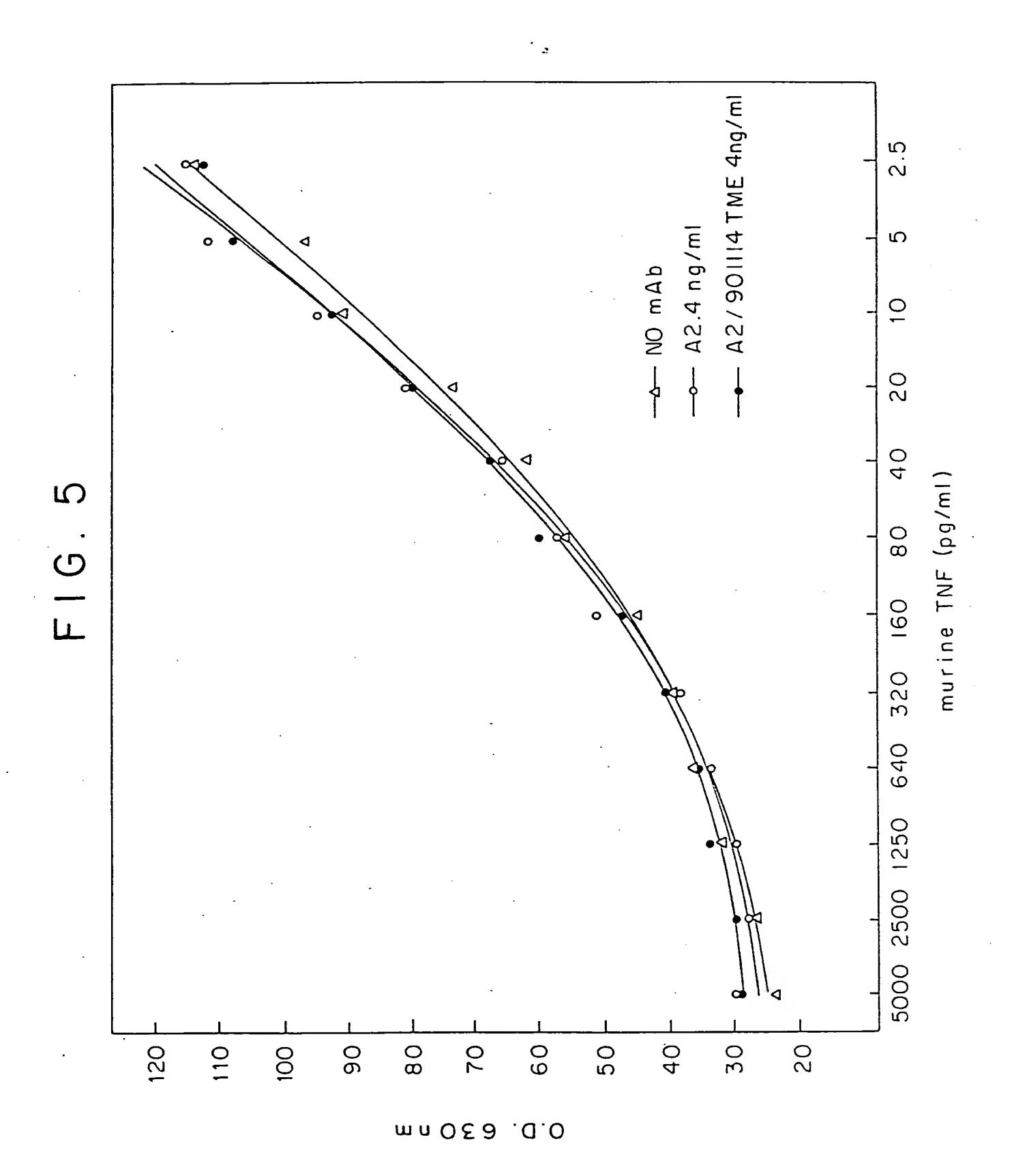


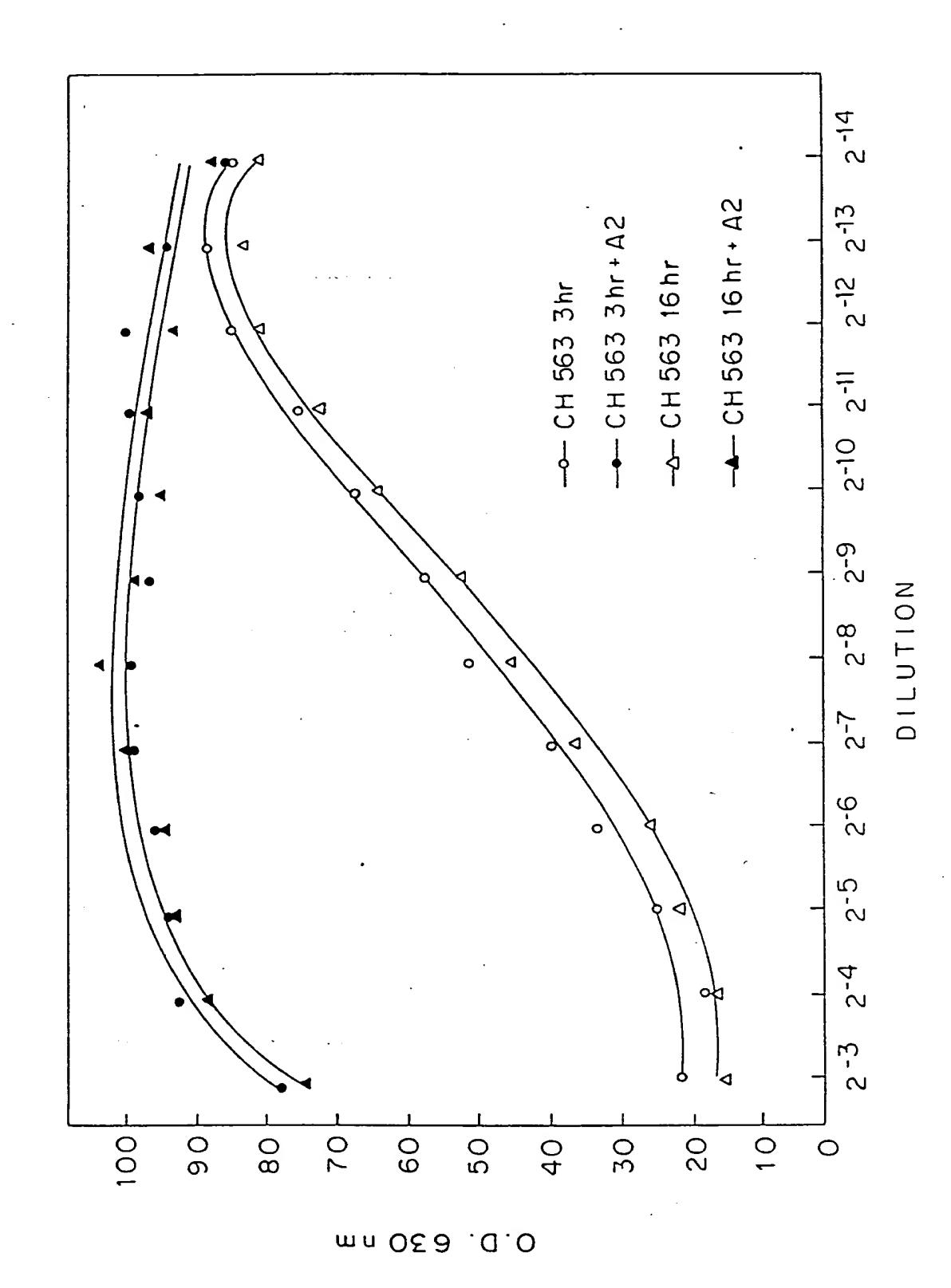
FIG. 3

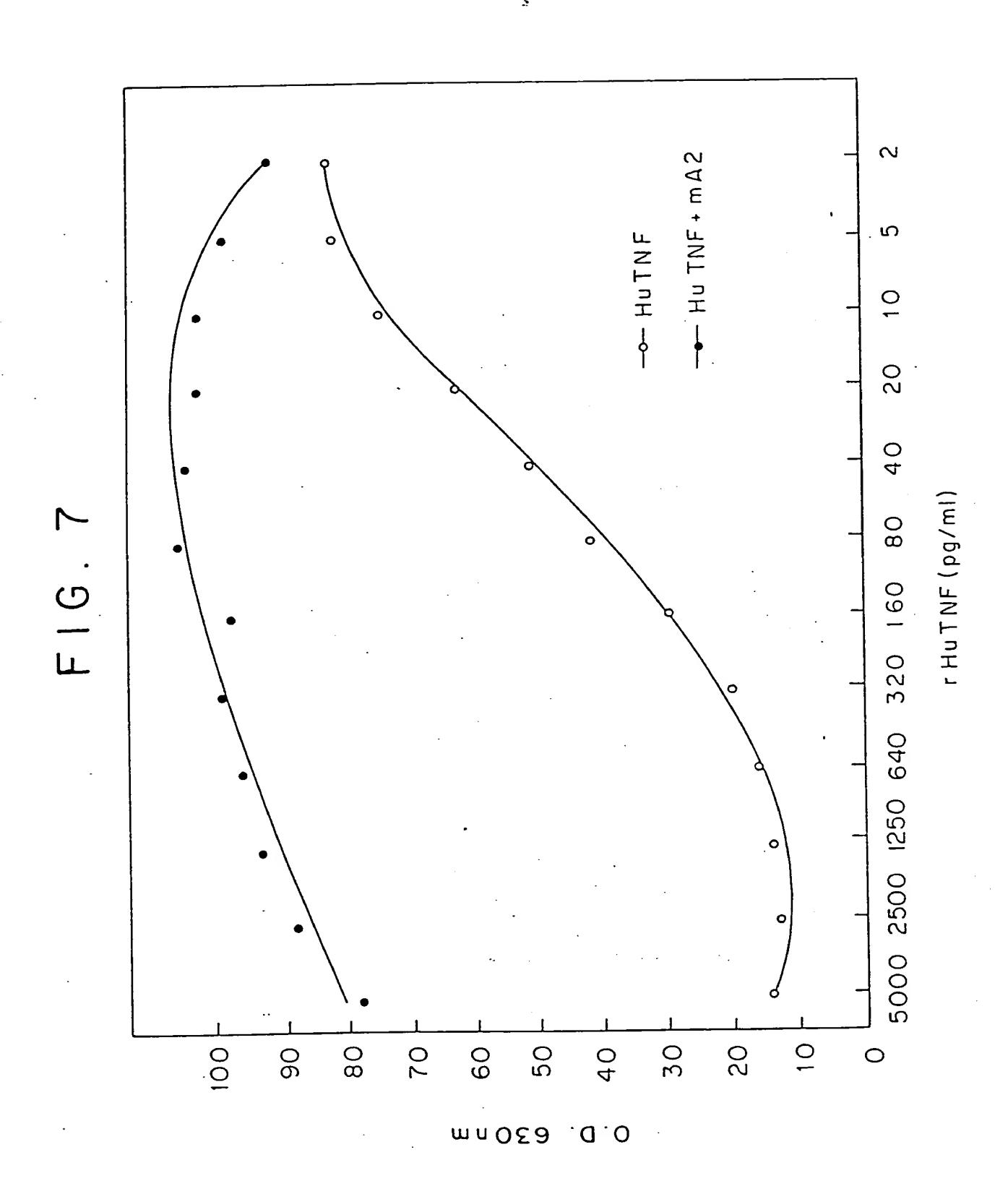




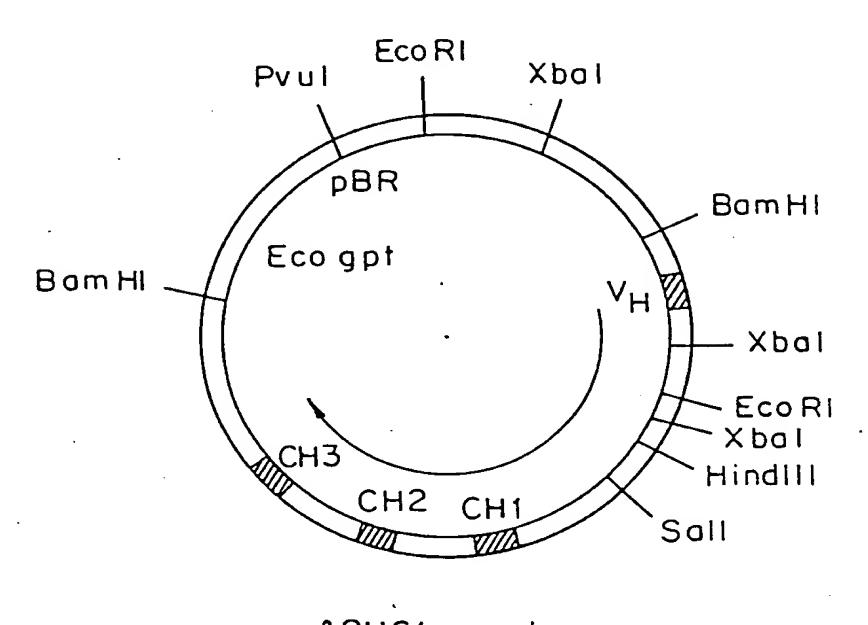


15



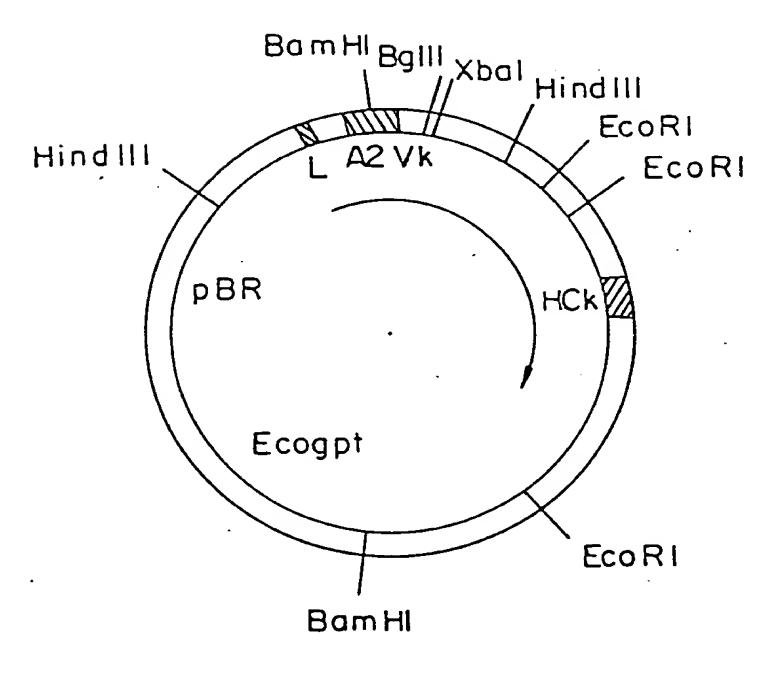


F 1 G. 8 a



pA2HG1apgpt

FIG.8b



pA2HuKapgpt

F I G. 9a

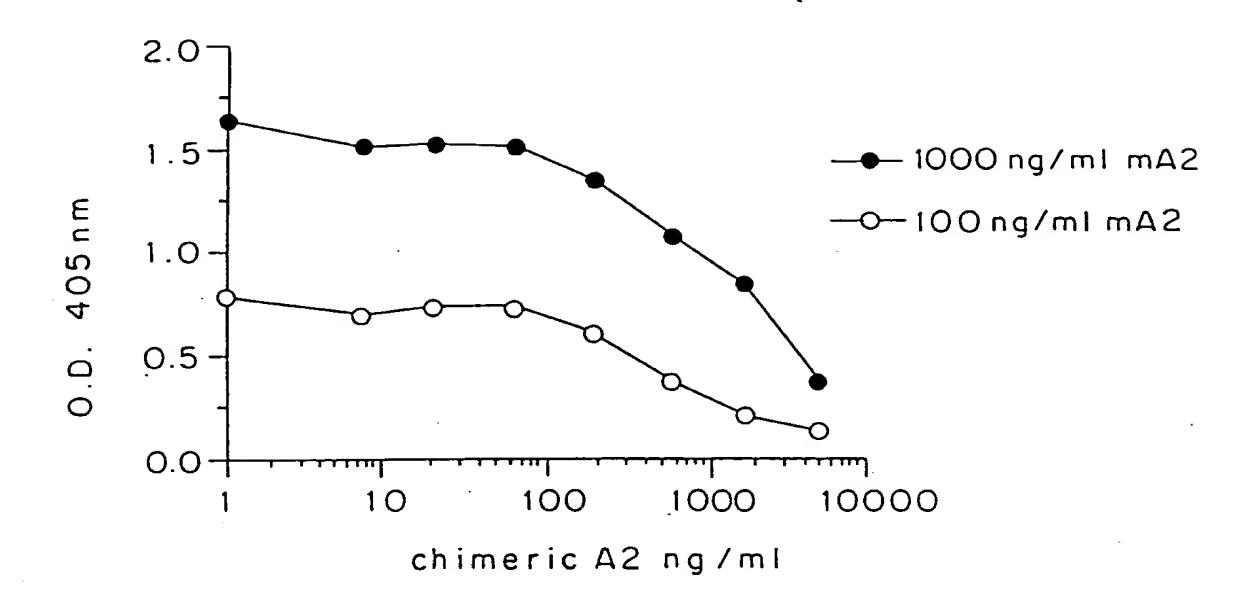
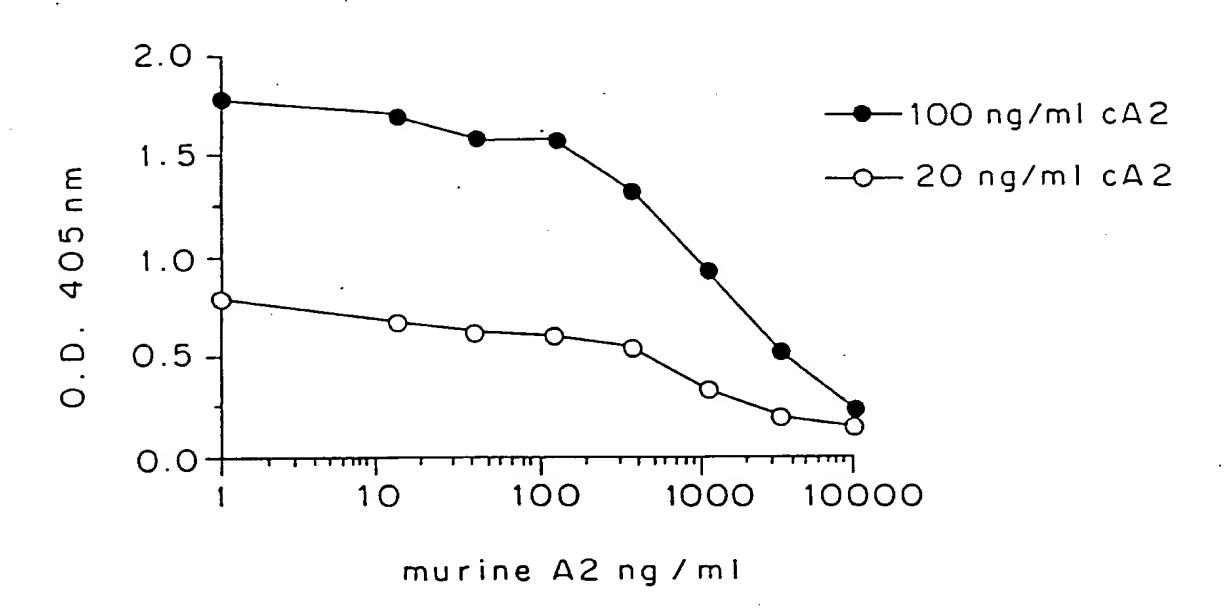


FIG. 9b



F I G. 10 a

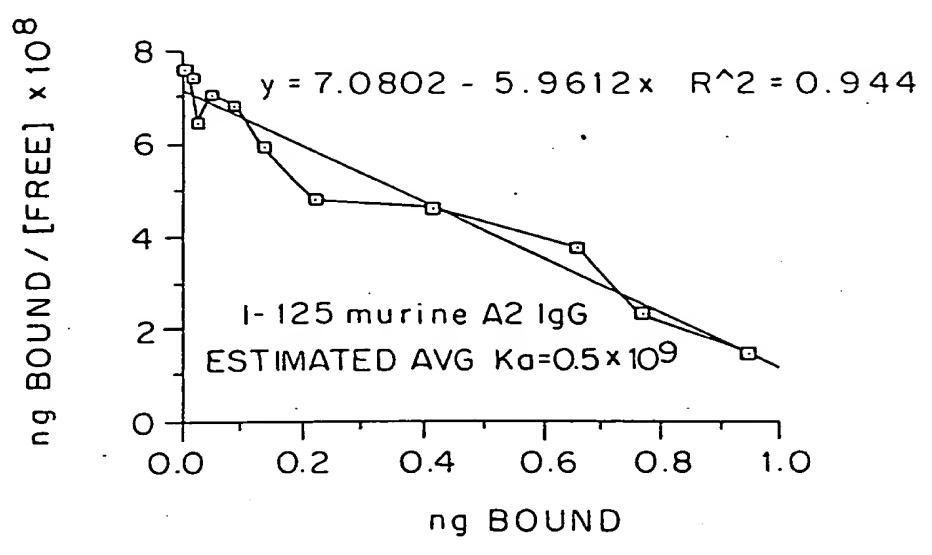
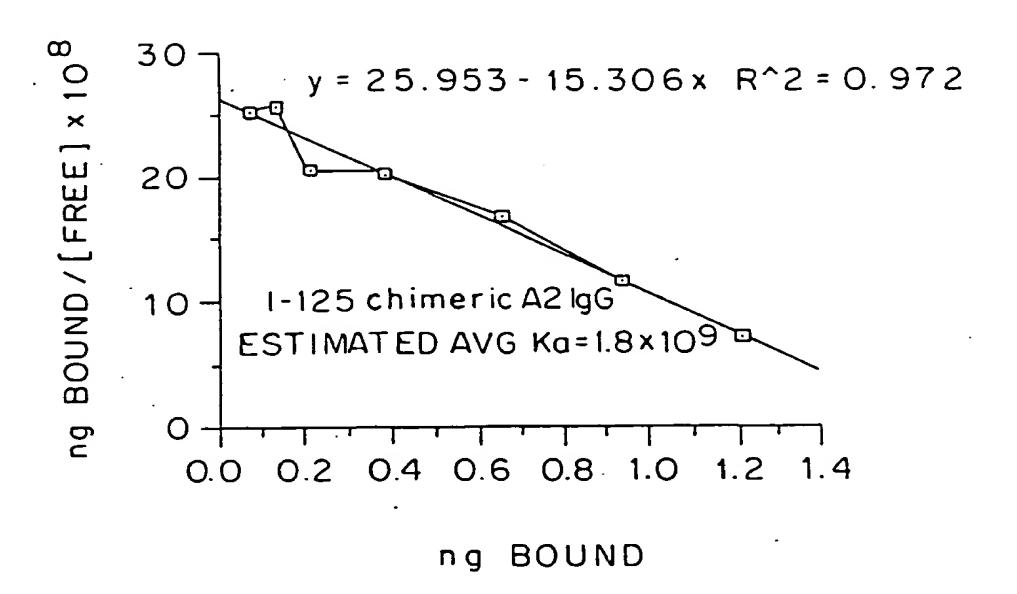
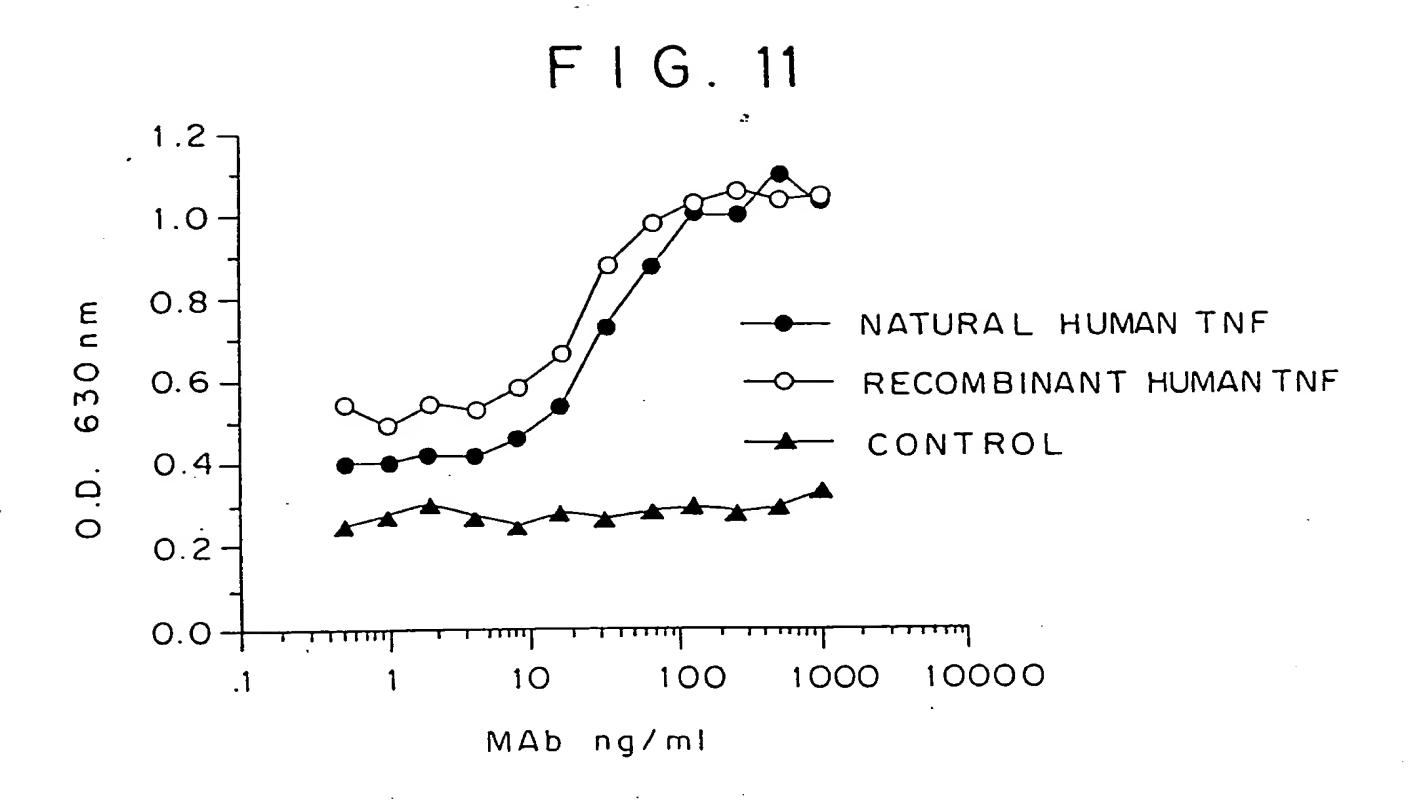
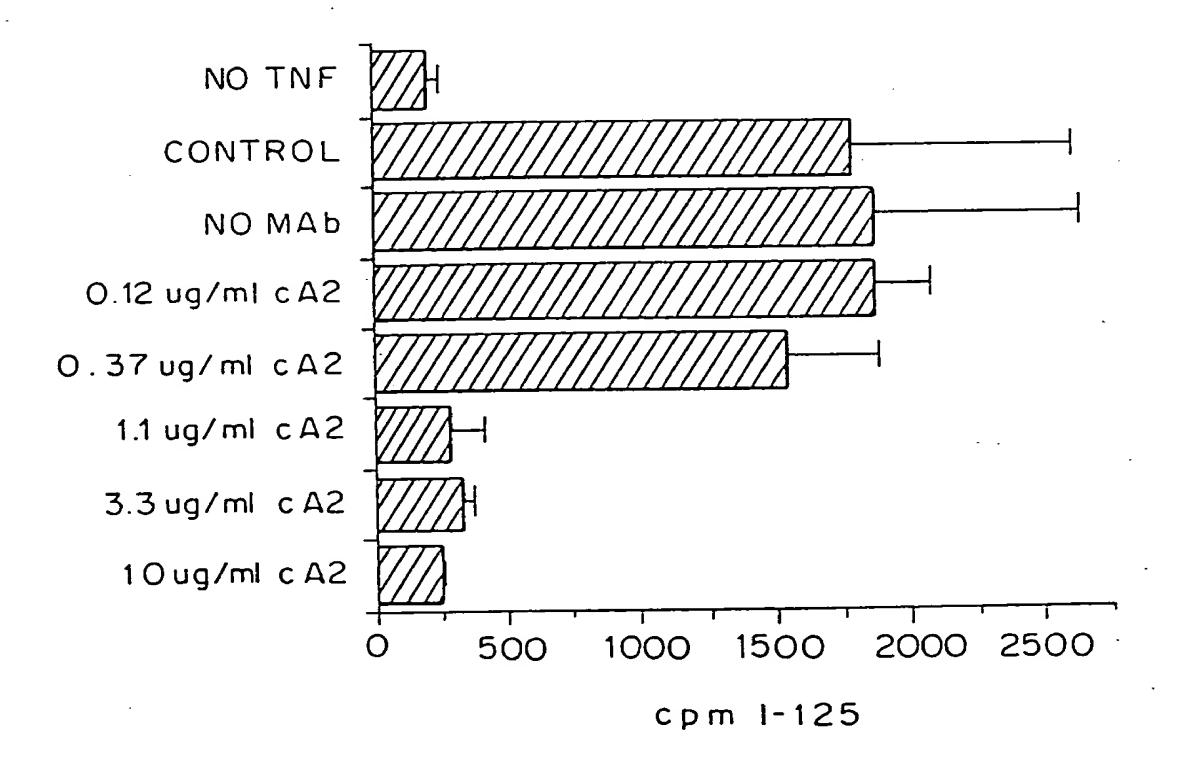


FIG. 10b





F I G. 12



F - G - 13

Pro Asn Ala Val Val His Ala Val Pro Lys 10 Asp Ser Pro Thr Arg Ser Ser Ser Arg Val

G1yAsn Ala Leu Leu Ala Asn Ala Arg Arg Asn 30 Leu Trp Gln Leu Gln GlyGlu Ala 21 Gln

Ser Tyr Ile Leu Tyr Leu G1yGlu Ser Pro 50 Val Val Leu Gln Asn Asp Arg Leu Glu Val 41

12 / Ile Thr His Thr Leu Leu Val His Thr Ser Pro 70 $\mathbf{C}\mathbf{y}\mathbf{s}$ Gly Gln Gly Lys Phe Leu Val Gln 61

4

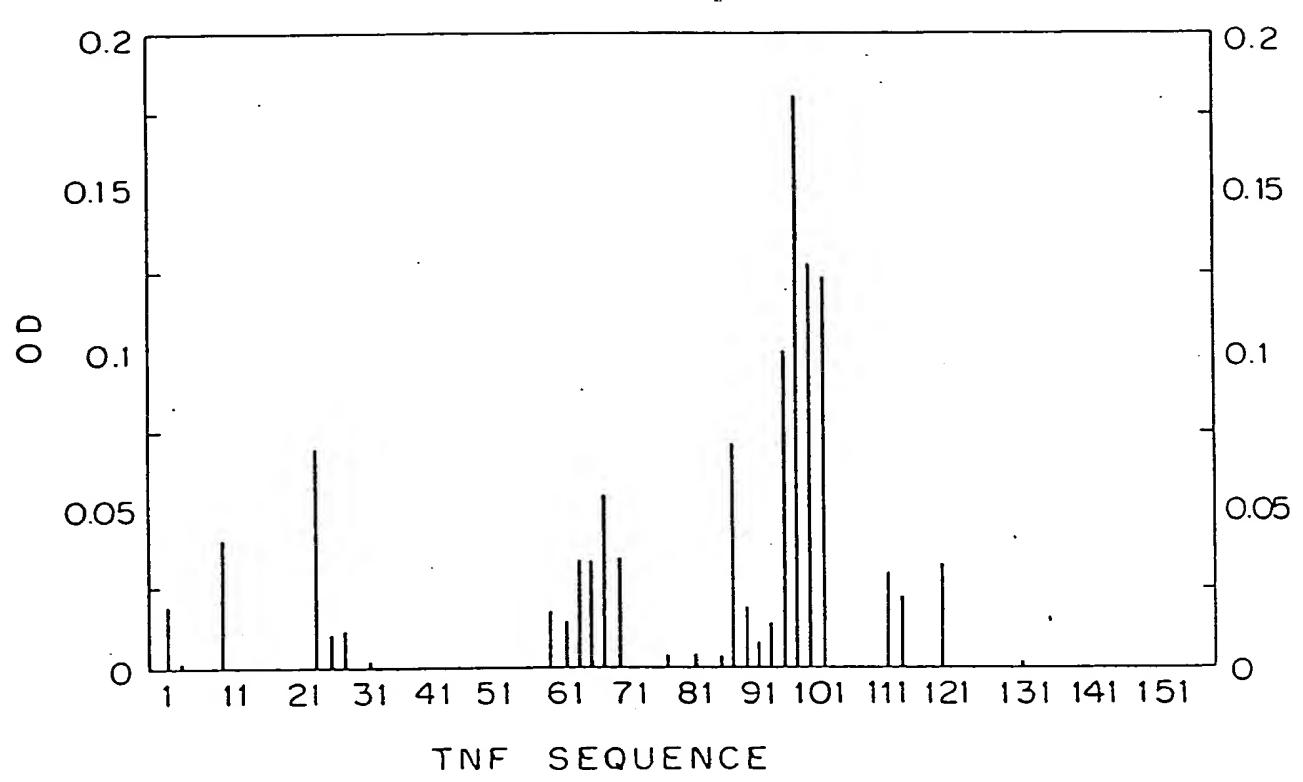
Pro Ser Lys Ile Ala Ser Leu Leu Asn Val 90Lys Gln Thr Tyr Ser Val Ala Ile Arg Ser 81

Leu TYr Ile Pro Glu Tyr Trp Pro Lys Ala 110 Glu Ala G1yGlu Pro Thr Glu Arg Gln Cys 101

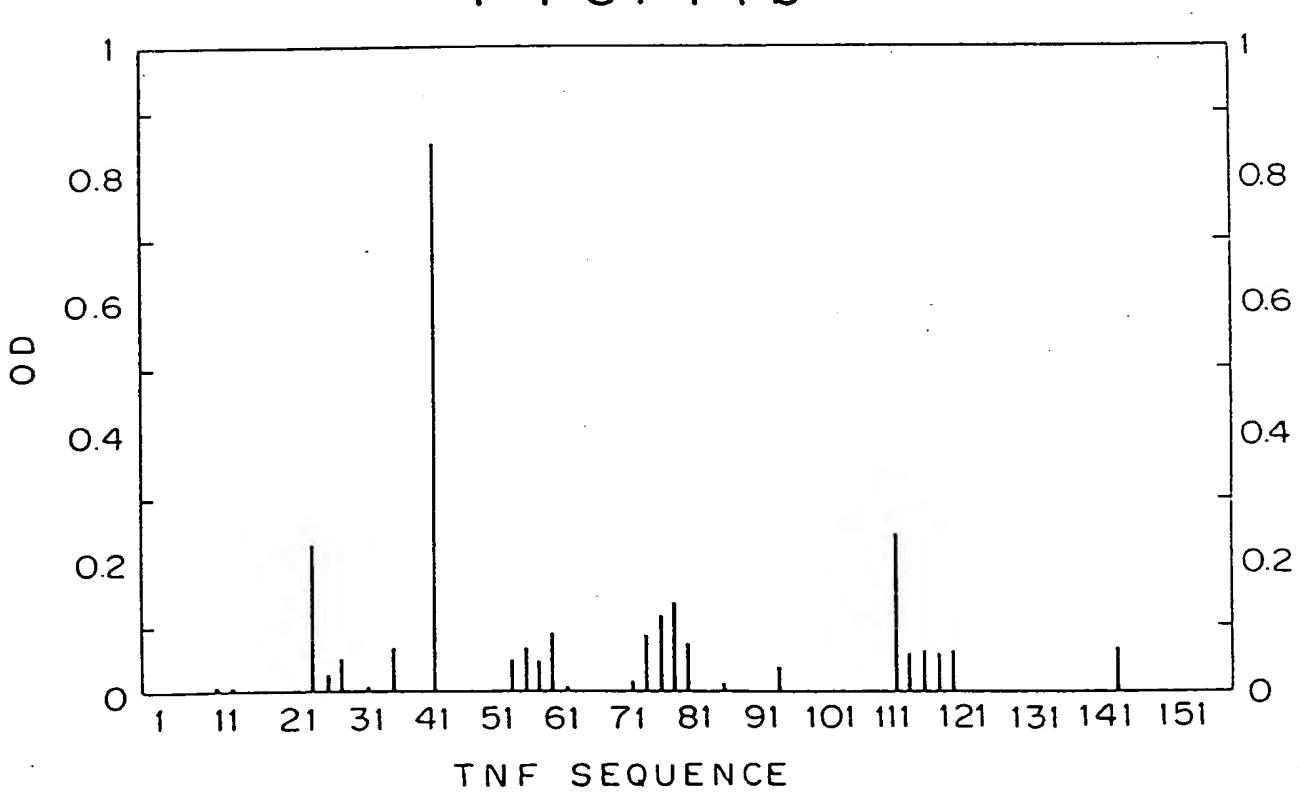
Asp Pro Arg Asn Ile Glu Ala Ser Leu Arg Asp 130 G1YLys Glu Leu Gln Phe Val G1y121 Gly

Leu Ala Ile Ile G1YPhe TYrVal 150 Gln G1ySer Glu Ala Phe Asp Leu Tyr 141





F I G. 14b



9

Pro Asn Ala Val Val His Ala Val Pro Lys Asp 10 Ser Pro Thr Arg Ser Ser Ser Arg 1 Val

G1yAsn Ala Leu Leu Ala Asn Ala Arg Arg Asn 30 Leu Trp Gln Leu Gln Gly Glu Ala 21 Gln

Ser TYr Ile Leu Tyr Leu G1YGlu Ser Pro Val 50 Val Leu Gln 41

Ile Thr ល Hi Thr Leu Leu Val His Thr Ser Pro 70 Cys Gly Gln Gly Asn Lys dsv Phe Arg Leu Leu Glu Val Gln Val 61

Pro H Se Lys Ile Ala Ser Leu Leu Asn Val 110 Glu Lys 90 Thr Gln Tyr Ser Val Ala Ile Arg Ser 81

1

Leu

TYr

Ile

Pro

Glu

Tyr

Trp

Pro

Lys

Ala

Ala

101

Asp Pro Arg Asn Ile Glu Ala Ser Leu Arg Asp 130 Lys Gly Gly Glu Glu Leu 0 Gln Thr Рће Gla Val Arg G1yGln 121 Gly Cys

Leu Ala Ile Ile G1YPhe TYr 150 Val Gln Gly Ser Glu Ala Phe Leu Asp 141 Tyr

Figure 16

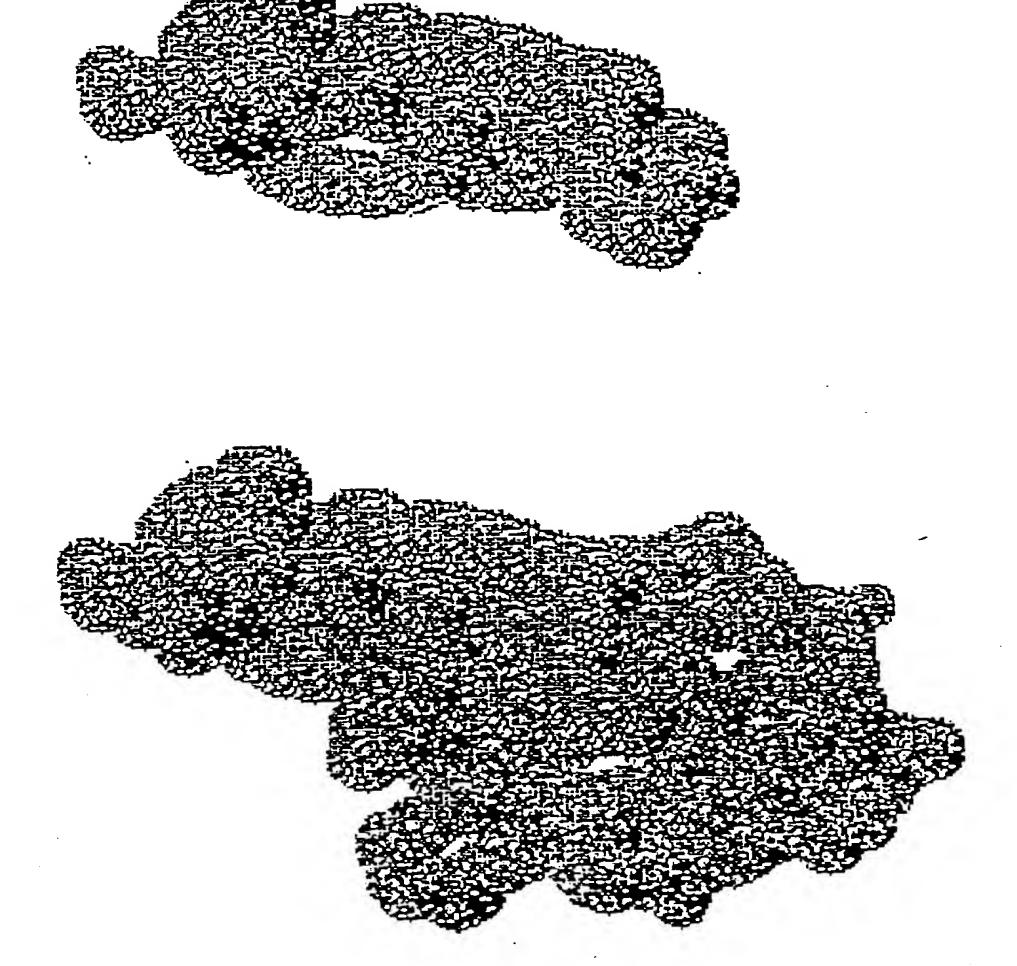


Figure 17A

CA2 LIGHT CHAIN VARIABLE REGION SEQUENCE

GACATCTTGCTGACTCAGTCTCCAGCCATCCTGTCTGTGAGTCCAGGAGAAAGAGTCAGT AsplleLeuLeuThrGlnSerProAlalleLeuSerValSerProGlyGluArgValSer

TTCTCCTCCACCCCAGTCAGTTCGTTGGCTCAAGCATCGACTGGTATCAGCAAAGAACA PheSerCysArgAlaSerGlnPheValGlySerSerIleHisTrpTyrGlnGlnArgThr

AATGGTTCTCCAAGGCTTCTCATAAAGTATGCTTCTGAGTCTATGTCTGGGATCCCTTCC AsnGlySerProArgLeuLeuIleLysTyrAlaSerGluSerMetSerGlyIleProSer

AGGTTTAGTGGCAGTGGATCAGGGACAGATTTTACTCTTAGCATCAACACTGTGCAGTCT ArgPheSerGlySerGlySerGlyThrAspPheThrLeuSerIleAsnThrValGluSer

GAAGATATTGCAGATTATTACTGTCAAGAAAGTCATAGCTGCCCATTCACGTTCGCCTCG GluAspIleAlaAspTyrTyrCysGlnGlnSerHisSerTrpProPheThrPheGlySer

GGGACAAATTTGGAAGTAAAA GlyThrAsnLeuGluValLys

∵∴ i

TCCTGTGTTGCCTCTGGATTCATTTTCAGTAACCACTGGATGAACTGGGTCCGCCAGTCT SerCysValAlaSerGlyPheIlePheSerAsnHisTrpHetAsnTrpVelArgGlnSer

CCAGAGAAGGGGCTTGAGTGGGTTGCTCAAATTACATCAAAATCTATTAATTCTGCAACA ProGluLysGlyLeuGluTrpValAlaGluIleArgSerLysSerIleAsnSerAlaThr

CATTATGCGGAGTCTGTGAAAGGGAGGTTCACCATCTCAAGAGATGATTCCAAAAGTGCT HisTyrAlaGluSerValLysGlyArgPheThrIleSerArgAspAspSerLysSerAla

GTCTACCTGCAAATGACCGACTTAAGAACTGAAGACACTGGCGTTTATTACTGTTCCAGG ValTyrLauGlnHctThrAspLauArgThrCluAspThrClyValTyrTyrCysSerArg

AATTACTACGGTAGTACCTACGACTACTGGGGGCCAACGCACCACTCTCACAGTCTCC AsnTyrTyrGlySerThrTyrAspTyrTrpGlyGlnGlyThrThrLeuThrValSer

Figure 17B

anti-TNF Trial

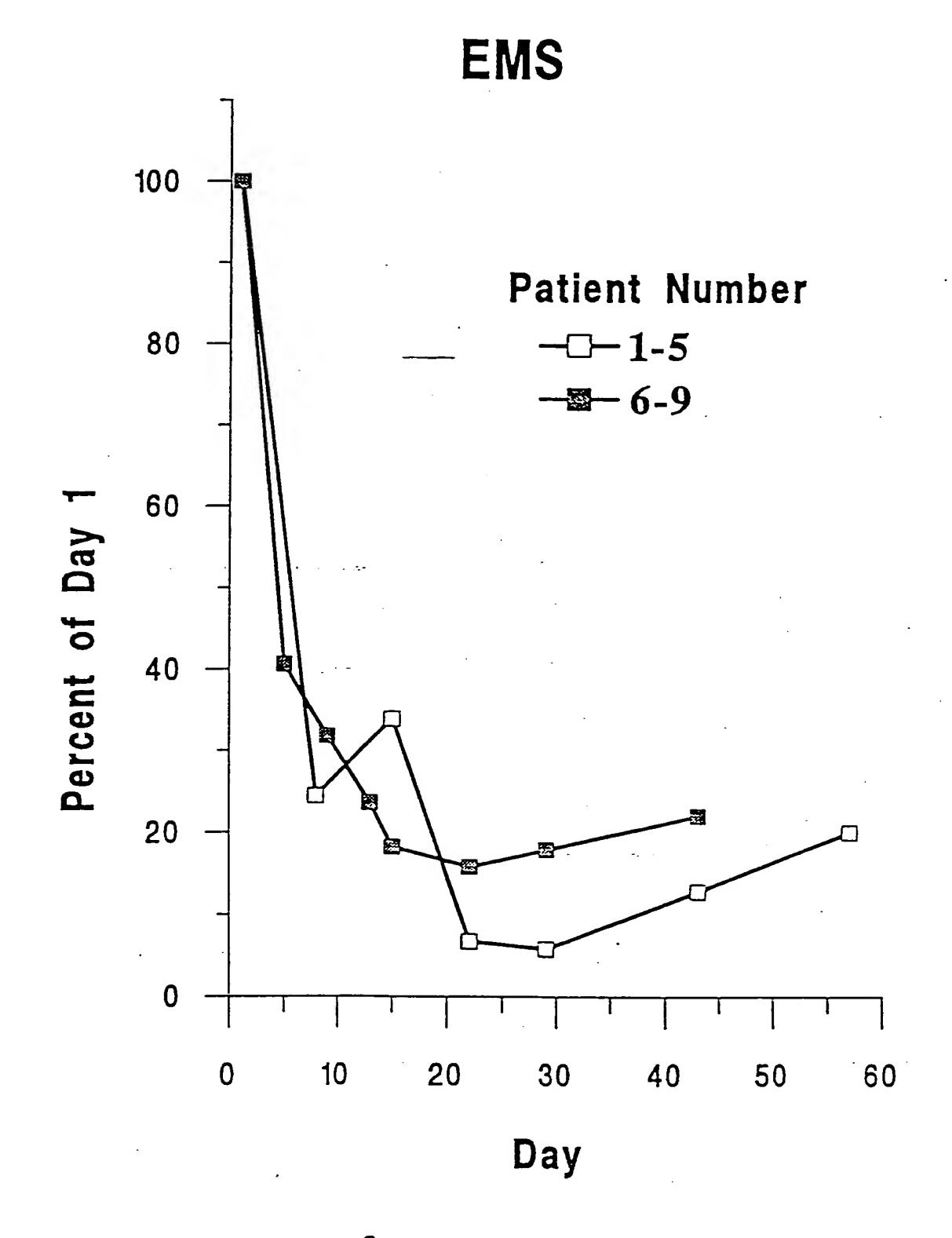
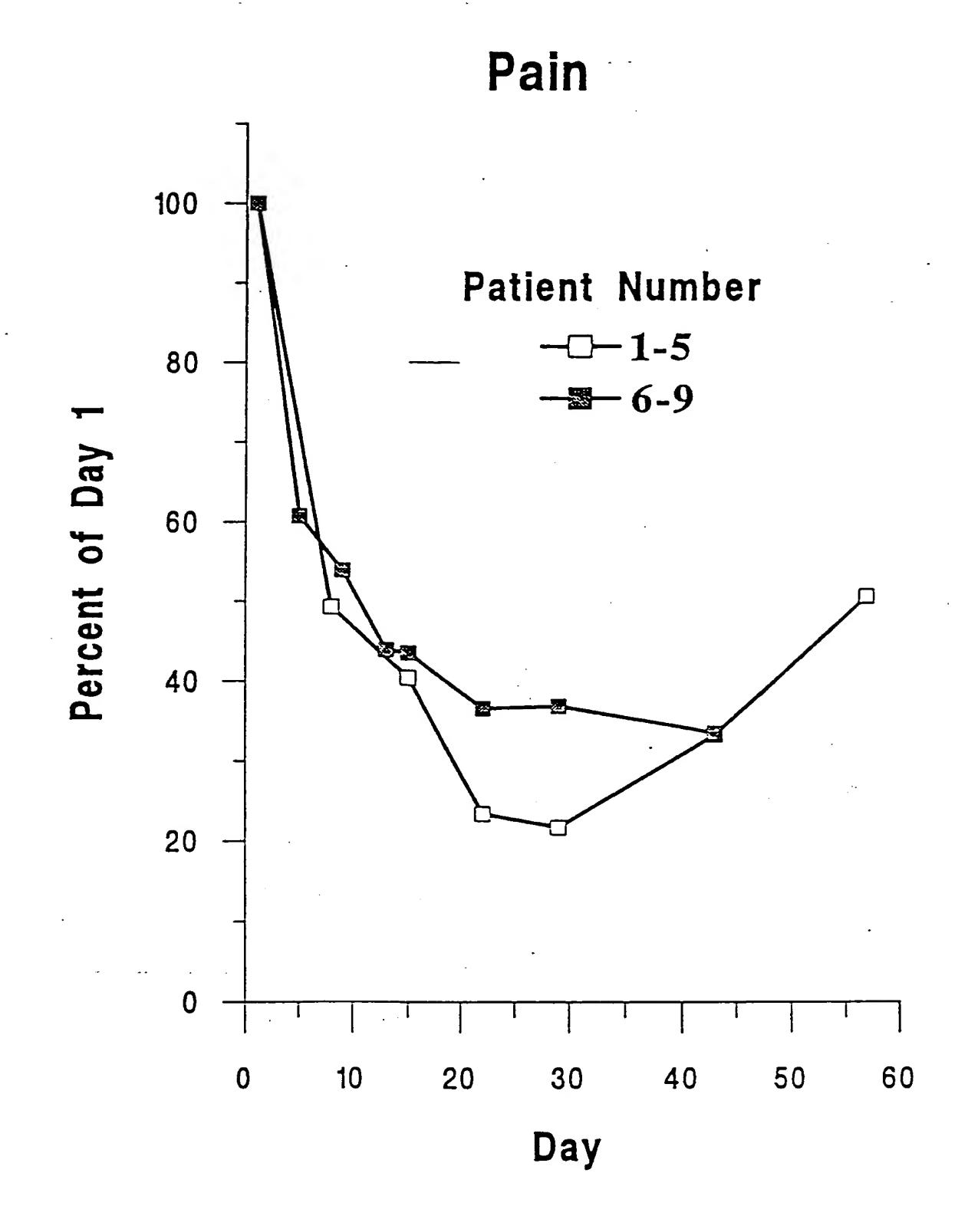


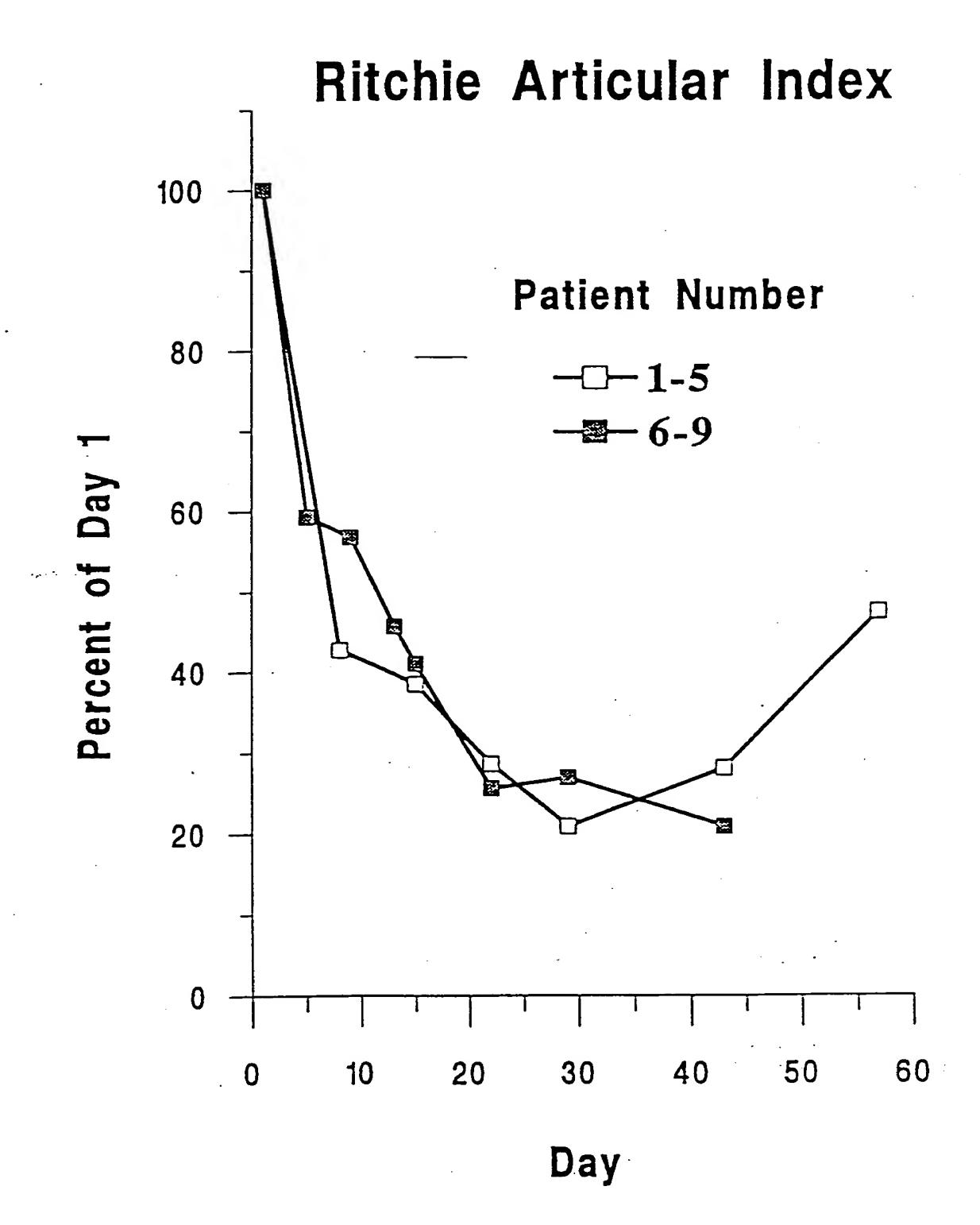
FIG-IS





F16.19

anti-TNF Trial



F16. 20

anti-TNF Trial

Swollen Joints

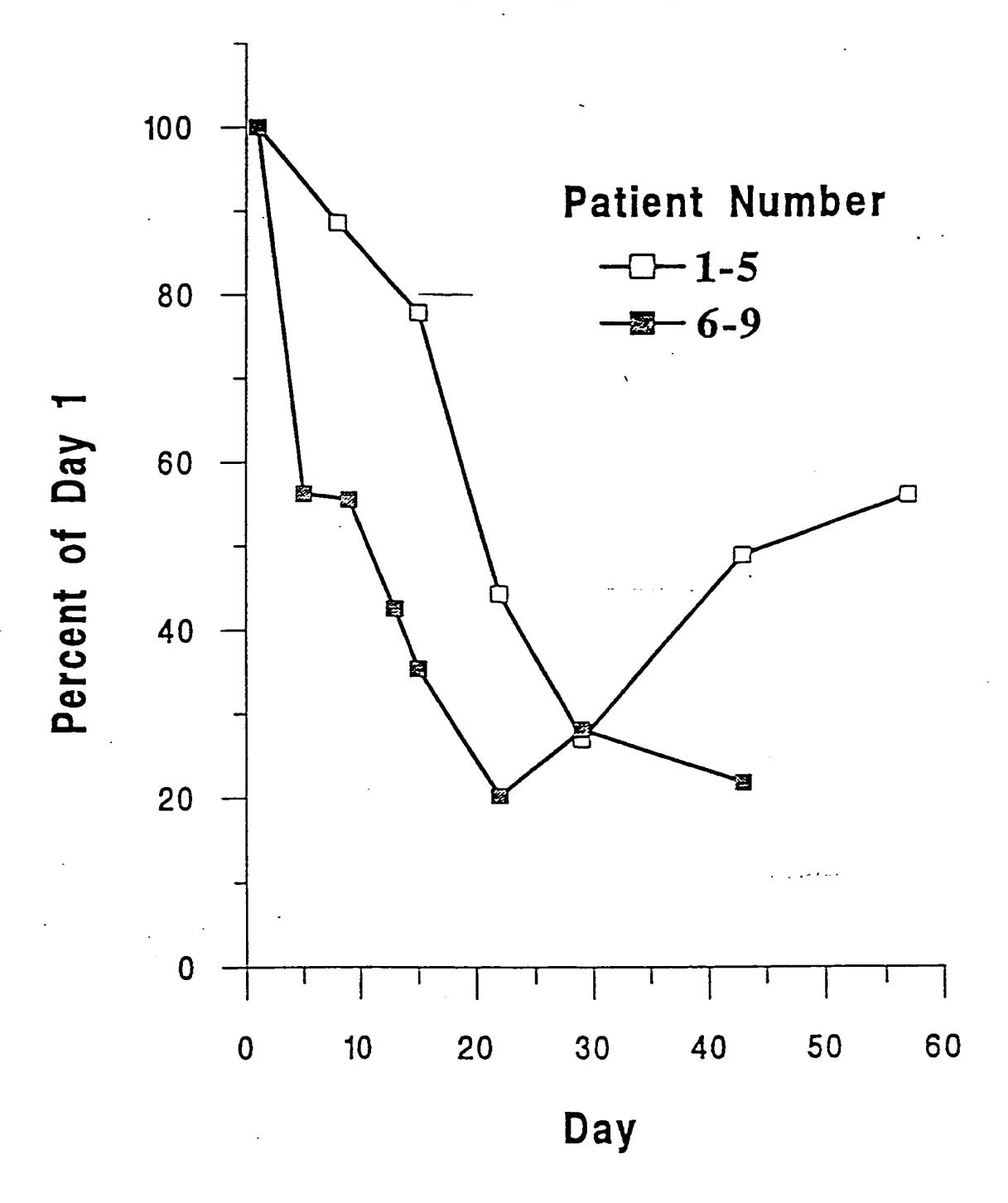
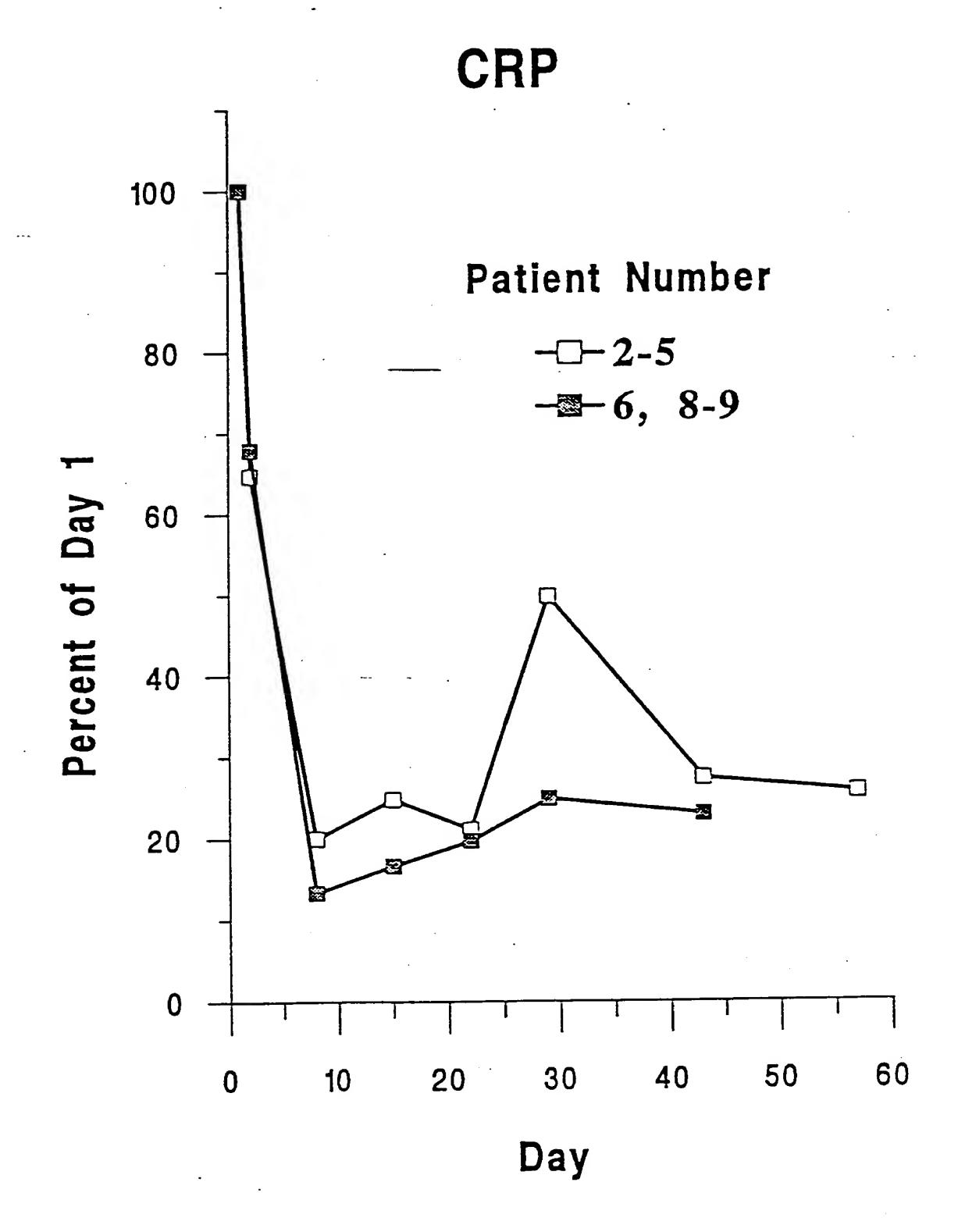


FIG 21





F16. 22



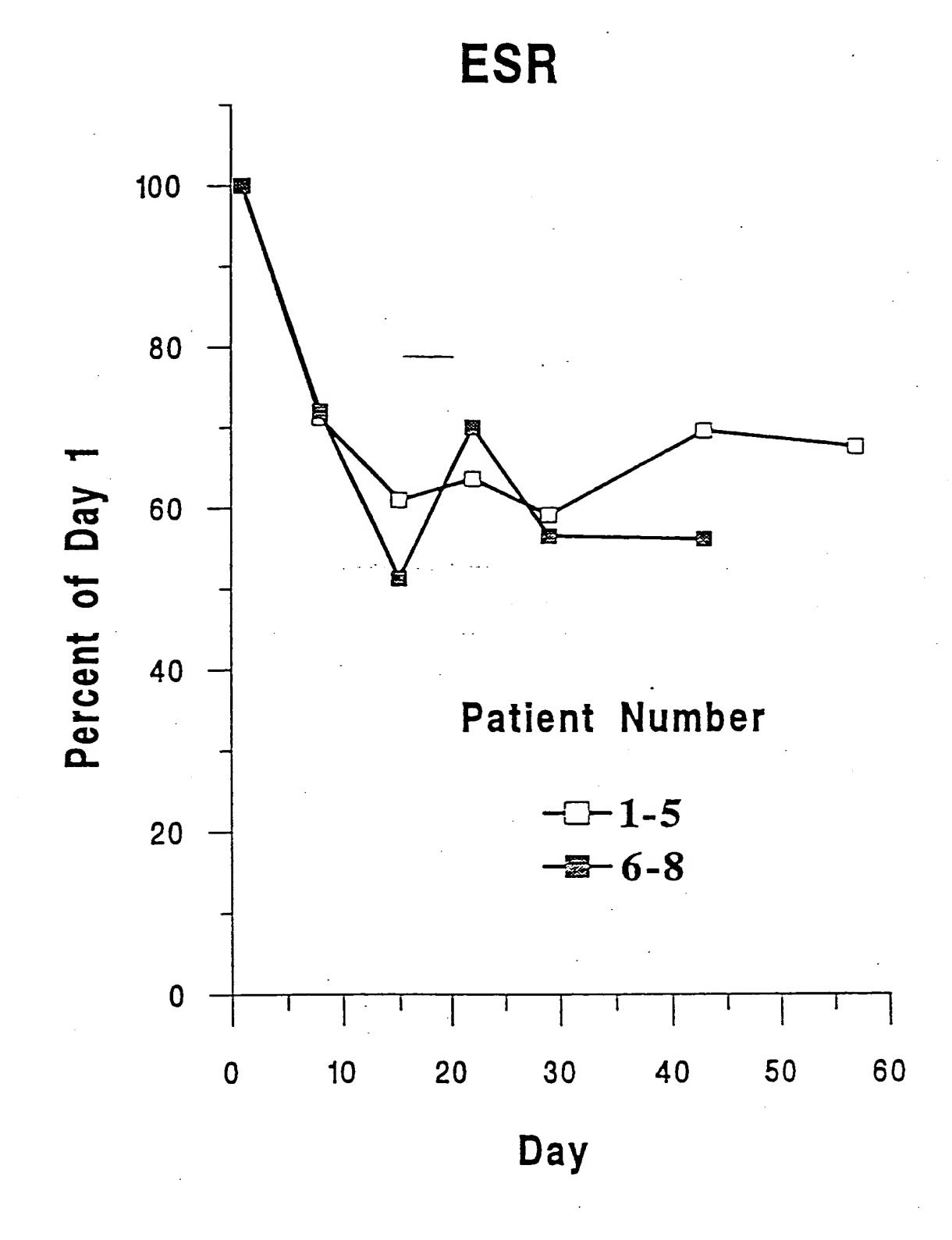


FIG 23



Index of Disease Activity

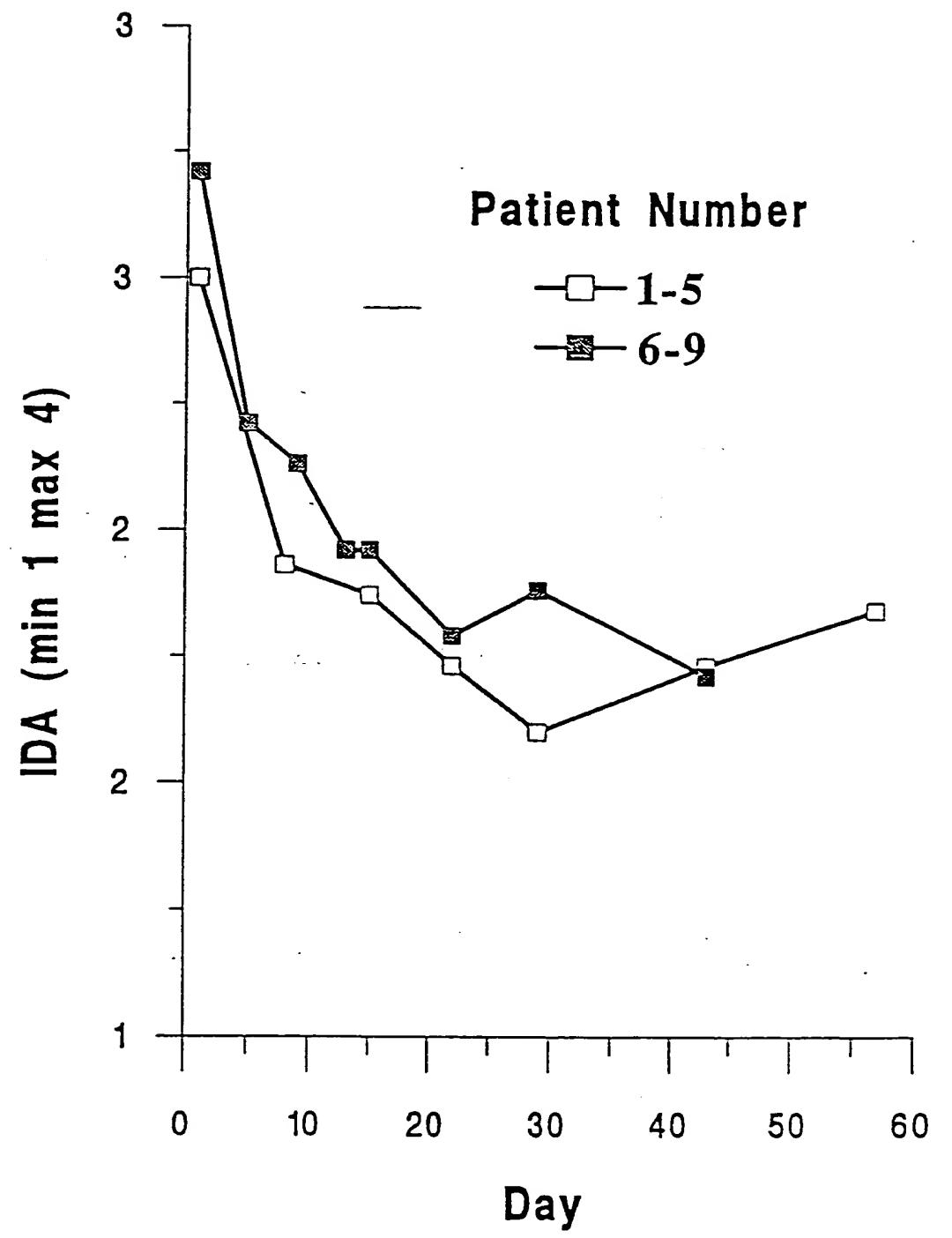


FIG 24